

Where Should I Run My Job?

Webinar Presentation, Dec. 7, 2011
NASA Advanced Supercomputing Division

Outline



- 1. Available Pleiades hardware (compute, front-ends, bridge nodes)
- 2. Running on pfe's and/or bridge nodes
- 3. Running on compute nodes
- 4. A sample PBS script
- 5. How to allow my job to run on any Pleiades processor type
- 6. Best practices with PBS jobs
- 7. PBS servers and queues

Supplemental Materials

- Job submission and monitoring
- Problems with submitting or running jobs

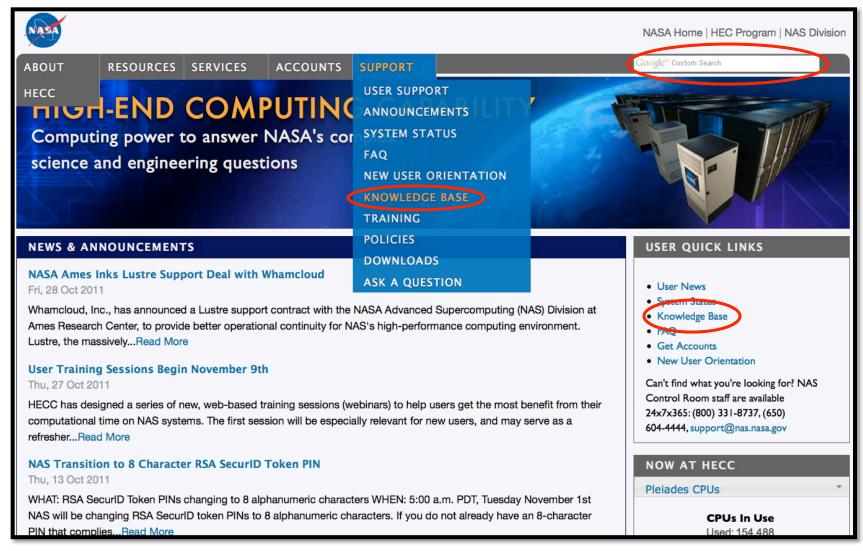


Preface: About NAS HECC Knowledge Base

- Most information in this webinar is available in Knowledge Base (KB)
- URLs to get to KB
 - NAS HECC home page: http://www.nas.nasa.gov/hecc/ then click on Knowledge Base
 - KB home page: http://www.nas.nasa.gov/hecc/support/kb/
- Searching information in KB
 - Each of the ~200 web pages (or articles) has an ID number
 - Can search with text or ID
- Relevant articles will be mentioned in this webinar using their IDs

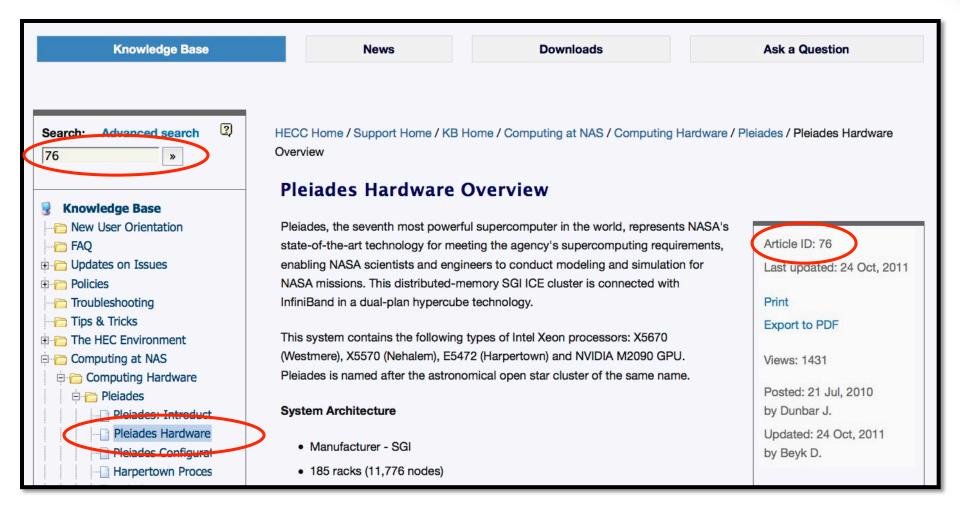
Searching info within HECC http://www.nas.nasa.gov/hecc/





Searching info within KB http://www.nas.nasa.gov/hecc/support/kb/





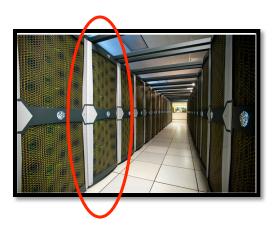
Available Pleiades hardware (compute, front-ends, bridge nodes)



➤ Compute nodes (KB 76 – 81)

- Hierarchy: \sim 185 racks -> \sim 185x64 nodes (r[1-222]i[0-3]n[0-15]) -> \sim 113,000 cores
- Three Intel Xeon processor types:

	Harpertown	Nehalem	Westmere
# of Racks	91	20	74
# of nodes	5,824	1,280	4,672
Cores/node	8	8	12
CPU speed	3.0 GHz	2.93 GHz	2.93/3.06 GHz
Memory/node	8 GB*	24 GB	24 GB
Memory/core	1 GB	3 GB	2 GB



1 rack has 64 nodes

* One Harpertown rack (rack 32) has 16 GB per node

- 64 Westmere nodes also include NVIDIA GPU (rack 219)
- Additional new hardware arriving Spring of 2012
- Compute nodes are for PBS jobs, not for logins

Available Pleiades hardware (compute, front-ends, bridge nodes) (cont.)



- Front-end pfe1-12 & bridge1-4 nodes (KB 76)
- Users can login to these nodes using ssh through sfe1-2 (see KB 59 for details)
- They are shared with other users for interactive processing
- Differences among them

	pfe1-12	bridge1-2	bridge3-4
processor	Harpertown	Harpertown	Nehalem-EX
cores/node	8	8	32
CPU speed	3.0 GHz	3.0 GHz	2.27GHz
Memory/node	16 GB	64 GB	256 GB
Network	1 GigE	10 GigE	10 GigE
os	SLES11SP1	SLES11SP1	SLES11SP1

Can run more processes on bridge[3,4]!!

bridge[1-4] have more memory and faster network than the pfe's !!

After installing new hardware in Spring of 2012, some nodes may use different
 OS version – be careful with compiling codes and checking for available software modules. To check for OS version, use the command:

uname –r or cat /etc/sgi-release

Running on pfe's and/or bridge nodes



Usage Guidelines (KB 181)

- What should run
 - Pfe's: compiling, short debugging, testing, jobs submission
 - Bridge: same
 - + pre/post-processing that need more memory
 - Tecplot no license restriction on number of users

Do not set stacksize to unlimited

IDL – 10 users (use "module load idl/8.1","Imstat –a" to check)

Matlab – 8 licenses (use "module load matlab/2010b", "matstat" to check)

- + file transfers to/from Lou/Columbia
- What cannot/should not run
 - MPI jobs (mpiexec is jailed)
 - A job that needs (n cores x m minutes) > 160
 - A job that needs more memory than maximum limit
 limits: pfe's: 8GB; bridge[1,2]: 56 GB; bridge[3,4]: 192 GB
- Recommend to check if there is enough free memory on the node first
 - Use the command top or cat /proc/meminfo to check

Running on the pfe's and/or bridge nodes (cont.)



- Do not over-utilize these nodes. Jobs are monitored.
 - CPU time usage monitored by "topGun" script
 - Memory usage monitored by "query_wms" script

Sample emails to user when overutilization is detected

From: topGun@nas.nasa.gov

Subject: interactive job limit exceeded on pfe9...

Date: July 29, 2011 7:53:33 AM PDT

To: jsmith@nas.nasa.gov

the following interactive job(s) on pfe9 have been found,

(but NOT killed).

Try fewer CPU's, less runtime -or- submit an interactive job session to PBS, via the command "qsub -I -I ncpus=1".

From: query_wms@nas.nasa.gov

Subject: [warning] jsmith@pfe1 over memory limit...

.. the following memory intensive job(s) found on pfe1

(but not killed)

```
USER PID RSS TIME COMMAND ismith 11690 10.5G 1:06 a.out
```

- * policy: up to 8.0G of 15.6G memory may be used
- * adjusted: 10.9G (136%) for jsmith's job 'a.out'
- * formula: at 0.0% load, tracking jobs using 1+ cpu's

try using less memory, re-run on bridge[1-4] -or- submit an interactive PBS session, via "qsub -I -I ncpus=1:model=wes".

Running on the compute nodes



- What should run on the compute nodes
 - All production jobs (MPI, OpenMP, hybrid, serial)
 - Some development jobs or large/long debugging jobs
- Memory and processes/threads constraints

limit	Harpertown	Nehalem	Westmere
processes/node or threads/node	8	8 x 2*	12 x 2*
Memory/node	7.5 GB	22.5 GB	22.5 GB

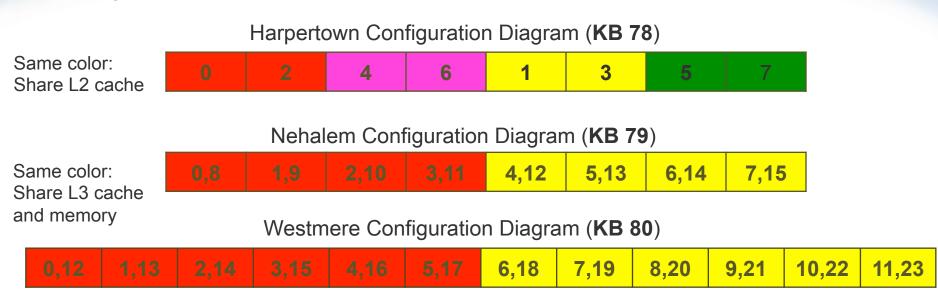
^{*}Due to hyper-threading; try with and w/o hyper-threading to see if there's benefit

- When node out-of-memory detected by PBS, email from support@nas.nasa.gov is
 sent to user. Try reduce the number of processes per node or use processor type(s)
 with more memory. See KB 216 for memory usage tips and memory monitoring tools
- The new hardware arriving in Spring will allow more memory/node (~30 GB/node)
- If you need large shared memory (~30 GB < mem < 4 TB) for production runs or have an OpenMP application that scales up to > 24 threads, apply for Columbia allocation

Running on the compute nodes (cont.)



Pinning processes/threads on the processors (here 1 processor means 1 core)



- CPUs that share identical resources (cache, memory, etc) are shown with the same color
- If processors are not fully occupied (e.g. running only 4 processes instead of 8 on Nehalem),
 different process placements will lead to performance differences. 4 processes assigned to
 0,1,2,3 on neh and wes, will likely cause resource contention and bad performance
- With mbind.x –cs, processes are assigned to cores in a spread format to minimize contention. So, 0,2,4,6 will be used on neh; 0,3,6,9 used on wes. –v prints pinning information to output.
- mbind.x works for MPI (SGI MPT, Intel MPI, MVAPICH, OpenMPI), OpenMP and hybrid
- More info available at: /u/scicon/tools/bin/mbind.txt

Running on the compute nodes (cont.)



- Cost for running a PBS job on the three process types
 - Resources are allocated and charged by nodes, not by cores
 - Running jobs do not share the same nodes
 - Projects are given SBUs each year for running PBS jobs
 - Use acct_query -pall -call -ujsmith -olow to get SBUs charged to jsmith's jobs today

Hypothetical example: running with 128 processes

	# nodes	Walltime (hrs)	SBU rate	Cost (SBU Hrs)
Harpertown	16	20	0.45	144
Nehalem	16	11	0.8	140.8
Westmere	11	12	1.0	132

- Which process type allows a faster turn-around?
 - Tools for checking how many free nodes for each processor type:
 - + qstat -au foo [@pbspl1] (or @pbspl3) []: means optional
 - + /u/scicon/tools/bin/qs [-s pbspl1] (or -s pbspl3)
 - + http://www.nas.nasa.gov/hecc/support/system_status.html (updated every 1-2 min)
 - Tools for checking how many nodes of each type queued jobs are waiting for:
 - + qstat i W o = + model, mission [@pbspl1] (or @pbspl3)
 - + /u/scicon/tools/bin/node_stats.sh

A sample PBS script



Sample PBS Script: (KB 175, 190)

```
resource list ->
                      #PBS –lselect=6:ncpus=8:mpiprocs=8:model=neh,walltime=24:00:00
                      ##PBS -lwalltime=24:00:00
     comment out ->
                     #PBS -q long
     queue name ->
  non-default GID ->
                      #PBS -W group list=s0101
                      #PBS -N my job name
  merge stdout/err ->
                     #PBS -j oe
       send email -> I
                     #PBS -m e
                     #PBS -r n
         no rerun ->
    load modules ->
                      module load comp-intel/11.1.072 mpi-sgi/mpt.2.04.10789
job submitted from ->
                     cd $PBS O WORKDIR
   implies –np 48 ->
                     mpiexec ./a.out > output
                      # or
  > recommended ->
                      # mpiexec -np 48 ./a.out > output
```

- Default GID can be found in /etc/passwd file (grep username /etc/passwd)
- PBS output/err file names: my_job_name.oxxxx (e.g., test.o12345, test.e12345)
 These files should be copied over by PBS to your \$PBS_O_WORKDIR when job ends
- PBS automatically restarts a job (up to 3 times) which fails due to system issues or if PBS fails to detect job running out of memory

How to allow my job to run on all three processor types?



Make sure that the executable can run on all three types

```
compile code without –xSSE4.2 (-xSSE4.2 runs on neh, wes only) or compile with -axSSE4.2,xSSE4.1 (KB 99)
```

Prepare three separate PBS scripts
 If not all cores are used, use mbind.x for pinning for good performance

```
For example, for a 12-process job, if each process needs 5 GB of memory #PBS –lselect=12:ncpus=1:model=har (with 7.5 GB, can fit only 1 process) #PBS –lselect=3:ncpus=4:model=neh (with 22.5 GB, can fit 4 processes) #PBS –lselect=3:ncpus=4:model=wes
```

```
#mpiexec –np 12 ./a.out > output
mpiexec –np 12 /u/scicon/tools/bin/mbind.x –cs -v ./a.out > output
```

recommend adding /u/scicon/tools/bin in your path in .cshrc set path = (\$path /u/scicon/tools/bin)

Best practices for PBS jobs



- Redirect application output to a file (KB 183)
 - mpiexec a.out > output (can monitoring job progress, since output will be in \$PBS_O_WORKDIR)
 - Without redirection, they are added to PBS stdout/stderr in head-node%/PBS/spool
 - /PBS/spool only has 1 GB of space. When it is filled up, PBS job may die. Won't get PBS stdout/err file.
 - New rule to be implemented soon: if PBS stdout/err > 100 MB, job is terminated
- Checkpoint long-running jobs
 - Checkpoint regularly to avoid losing data when job runs out of wall-time or fails due to system issues
 - If not sure how long it takes to finish the run, can checkpoint prior to running out of wall-time
 (KB 199, pbs_time_left.a)
 - Turn on #PBS –r n if you don't want PBS to restart your job for you
- Avoid combining large-cpu number crunching with small-cpu post-processing in 1 PBS job
 - If the small-cpu post-processing takes a long time, resource and allocation are wasted.
 - Submit a separate PBS job with fewer nodes for post-processing or do it on bridge nodes if possible
- Package multiple serial runs in a single PBS job (KB 184)
 - Submitting too many jobs slows down PBS job scheduling
 - Package serial jobs with module load mpi-mvapich2/1.4.1/intel and use technique described in KB 184

PBS servers and queues



- > PBS (KB 126, 173-179, 186-190, 290)
- Two PBS servers:
 - pbspl1 is the default, it manages most queues
 - pbspl3 manages 2 queues (devel & gpu) (KB 290)
- Some queues (normal, long, debug, devel, gpu) are accessible to all Pleiades users
- Some are accessible only to certain missions/groups/users
 For example, armd_spl, kepler, vlong (16 days max)
 Manager(s) approval are needed to create/access these queues
- Reservation queues (KB 176) require HECC manager approval; set up by sysadm
 Special circumstances which warrant the use of reservation queues:
 - time critical simulations over a period of time
 - very large simulations (e.g., 20,000-core jobs)

Charged even when not used

PBS servers and queues (cont.)



> PBS queues accessible to all Pleiades users: (KB 187, 290)

```
pfe1% qstat -Q (or qstat -Q @pbsp11)
                             pfe1% qstat -Q @pbsp13
      Ncpus/
                Time/
Queue
                             Oueue
                                    Ncpus/
                                            Time/
        max/def max/def
                                     max/def max/def
name
                          pr
                             name
                                                      pr
==
debug
       1025/
             8 02:00/ 00:30
                          15
                             devel
                                    4800/ 1 02:00/
long 8192/ 8 120:00/ 01:00
                           0
                             gpu
                                      --/ 8 08:00/01:00
                                                      0
normal
             8 08:00/ 01:00
```

- Debug queue has higher priority
- If queue name is not specified during job submission:
 walltime <= 8 hours, default to normal queue
 walltime > 8 hours, but <=120 hours, long queue
 walltime > 120 hours, job will be rejected
- har, neh, wes available for all these queues
- if processor type not specified, default to harpertown

- 8 Westmere racks (512 nodes) set aside
 7x24 for devel queue
- Served by pbspl3, devel queue allows faster job scheduling and turnaround
- 1 running job per user on devel queue
- no production work on devel queue
- Use gpu queue (64 nodes) if you have a CUDA application for running on GPU
- Use qstat -fQ queue_name@server_name to get more info about each queue

Summary



- Where should I run my jobs?
- Front-end and bridge nodes resources and usage guidelines
- Compute nodes resources and factors to consider for PBS jobs

•	three processor types	
•	cores consideration	

•	memory	consideration
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•	cost	consideration
	COSL	CONSIDERATION

Harpertown	Nehalem	Westmere
8	8x2*	12x2*
7.5 GB	22.5 GB	22.5 GB
SBU 0.45	0.8	1.0

(acct_query)

- availability consideration (qstat –au foo, qs, qstat –i, node_stats.sh)
- pinning for better performance (mbind.x cs v)
- Sample PBS scripts and Best Practices for PBS jobs (redirect output !!)
- PBS queues to submit your jobs to (qstat –Q, use devel for development)
- > For more information, see KB articles listed in this webinar
- For assistance: call 1-800-331-8737 or 650-604-4444 or email support@nas.nasa.gov



Slides Prepared by Sherry Chang

pdf and recording of this webinar will be available shortly at: http://www.nas.nasa.gov/hecc/support/training.html

Next webinar "I/O: Tips and Techniques" tentatively scheduled on Jan. 11, 2012

Suggestions for future webinar topics are welcomed

Job Submission and Monitoring



Submitting your PBS job

Batch jobs

```
% qsub job_script (for pbspl1 only) or %qsub –q devel@pbspl3 job_script 12345.pbspl1.nas.nasa.gov 2468.pbspl3.nas.nasa.gov
```

Options in the command line override those in job_script
 % qsub -lselect=4:ncpus=12:model=wes job_script

```
12347.pbspl1.nas.nasa.gov
```

Interactive PBS job (good for debugging or development work)

```
% qsub -I -lselect=4:ncpus=12:model=wes (Note: no job script should be included here)
```

Monitoring your PBS job

```
% gstat –u įsmith
                                                       (show all running/queued jobs by jsmith - pbspl1)
% qstat @pbspl1 @pbspl3 –W combine –u jsmith
                                                       (show all jsmith's jobs served by pbspl1 and pbspl3)
% qstat –nu jsmith
                                                       (show nodes for running jobs - pbspl1)
% qstat -f 2468.pbspl3
                                                       (show details of a running/queued job - pbspl3)
% qstat -s 12347
                                                       (gives one line status explanation of a job - pbspl1)
% qstat –i
                                                       (show all queued jobs sorted by priority – pbspl1)
% qstat –xu jsmith
                                                        (show all finished/running/queued jobs of jsmith)
% qstat –xf 12345
                                                       (show details of a finished/running/queued job)
```

Problems with submitting or running jobs



Can't Submit: (KB 197)

- qsub: Job exceeds queue and/or server resource limits
 (for example, asking for 3 hours in devel or debug queue)
- qsub: Job rejected by all possible destinations
 (for example, qsub –lwalltime=140:00:00)
- qsub: Unauthorized Request
 (AUID/GID not in ACL, no more allocation)
- qsub: Bad GID for job execution
 (AUID not in the GID used for submission)
- qsub: would exceed complex's per-user limit (exceed 300 total jobs in Q or R state)

Job waiting in queue, not running (KB 198)

- get a one line status information (use *qstat -s jobid*)
- job is waiting for resources (use *qstat -i* to check job position)
- job will never run due to impossible resource request (eq. submit to devel asking for har)
- job would exceed mission CPU share (use %qstat –W shares=- to check, KB 168)
- system going into dedicated time (check email and MOTD announcement)
- your home filesystem or /nobackup is down (possible system problems)
- over quota on home or /nobackup filesystem (hard limit)
- job on Hold (for example, held by sysadm or after being run 3 times but failed)

Problems with running jobs (cont.)



- What to do if my job ran, but failed or stalled?
- check PBS stdout/stderr (in \$PBS_O_WORKDIR) for possible error messages

command not found (check if module loaded? path included? hidden character?)
Disk quota exceeded (delete files to go under disk/inode quota hard limit)

```
SIGFPE(8) – Floating-point exception
SIGSEGV(11) – Segmentation violation
(recompile code with debugging options –O0 –g –check –traceback)
```

- check email if PBS detects job encountering OOM
 (monitor job to find out memory needed; increase memory request) (KB 216)
- possible system problems (such as IB network, Lustre file-system, bad cpus or memory)
 MPI: rank 100: r1i0n3 IB board mlx4_0 port 1 had fault
 (report problems; resubmit to different processor type)
 (the systems team adds check in the PBS prologue to offline low-memory nodes)
- ask for help (send email to support@nas.nasa.gov)